

PC I/US 97/11555

Patent document			7/03/3//11555	
cited in search report	Publication date	Patent tamily member(s)	Publication date	
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US 5160702 A	03-11-92	NONE		
WO 9708556 A	06-03-97	AU 6913196 A	19-03-97	
WO 9607919 A	14-03-96	US 5627041 A AU 3462795 A CA 2198854 A EP 0778950 A	06-05-97 27-03-96 14-03-96 18-06-97	



## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification 7: C12Q 1/68, B01L 3/00

A1

(11) International Publication Number:

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WO 00/40750

JEEEL1

(43) International Publication Date:

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PCT/EP99/10347

(22) International Filing Date:

23 December 1999 (23.12.99)

(30) Priority Data:

9828785.7

30 December 1998 (30.12.98) GB

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(72) Inventors; and

- (75) Inventors/Applicants (for US only): ORLEFORS, Anna, Edman [SE/SE]; Vaderkvarnsgatan 35B, S-753 26 Uppsala (SE). ERICKSON, Kerstin [SE/SE]; DagHammarskjoldsv 245 B, S-756 52 Uppsala (SE). LOFMAN, Esfir [SE/SE]; Borjegat 1B, S-753 13 Uppsala (SE). ANDERSSON. Per [SE/SE]; Hornsgatan 147, S-117 30 Stockholm (SE). ULFENDAHL, Per, Johan [SE/SE]; Rapphonsvagen 10B, S-756 53 Uppsala (SE).
- (74) Agent: ROLLINS, Anthony, John; Nycomed Amersham plc, Amersham Laboratories, White Lion Road, Amersham, Bucks HP7 9LL (GB).

(81) Designated States: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZA, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

### **Published**

With international search report.

(54) Title: METHOD FOR SEQUENCING DNA USING A MICROFLUIDIC DEVICE

### (57) Abstract

The present invention describes a method for sequencing DNA in a device with microfluidics properties and a set of reagents for its use. This microfluidic device may be in form of a disc with radially extending microchannel structures (CD form) having an inner application area that may be common for one or more microchannel structures. By spinning the disc the liquid can be driven from an applicator area into reaction and/or detection are as closer to the periphery of the disc. Liquid transportation may thus be driven by centripetal force. The microfluidic device may also have other gemoetrical forms. Several methods can be used to determine the sequence of DNA according to the invention but the real time determination of released pyrophosphate using the luciferase luciferin reaction is preferred.

### PATENT COOPERATION TREATY

From the INTERNATIONAL PRELIMINARY EX	AMINING AUTHORITY		
To:  ROLLINS, Anthony J.  NYCOMED AMERSHAM PLC Amersham Laboratories White Lion Road Amersham Buckinghamshire HP7 9LL GRANDE BRETAGNE		PCT  NOTIFICATION OF TRANSMITTAL OF THE INTERNATIONAL PRELIMINARY EXAMINATION REPORT (PCT Rule 71.1)  Date of mailing (day/montr/year) 11.01.2001	
Applicant's or agent's file reference			
PU9844			PORTANT NOTIFICATION
International application No. PCT/EP99/10347	International filing date (da 23/12/1999	ny/month/year)	Priority date (day/month/year) 30/12/1998

- 1. The applicant is hereby notified that this International Preliminary Examining Authority transmits herewith the international preliminary examination report and its annexes, if any, established on the international application.
- 2. A copy of the report and its annexes, if any, is being transmitted to the International Bureau for communication to all the elected Offices.
- 3. Where required by any of the elected Offices, the International Bureau will prepare an English translation of the report (but not of any annexes) and will transmit such translation to those Offices.

#### 4. REMINDER

The applicant must enter the national phase before each elected Office by performing certain acts (filling translations and paying national fees) within 30 months from the priority date (or later in some Offices) (Article 39(1)) (see also the reminder sent by the International Bureau with Form PCT/IB/301).

Where a translation of the international application must be furnished to an elected Office, that translation must contain a translation of any annexes to the international preliminary examination report. It is the applicant's responsibility to prepare and furnish such translation directly to each elected Office concerned.

For further details on the applicable time limits and requirements of the elected Offices, see Volume II of the PCT Applicant's Guide.

Name and mailing address of the IPEA/

Fax: +49 89 2399 - 4465

Authorized officer

European Patent Office

D-80298 Munich Tel. +49 89 2399 - 0 Tx: 523656 epmu d

AMERSHAM PHARMACIA BIOTECH AB et al

Danti, B

Tel.+49 89 2399-8161



### From the INTERNATIONAL BUREAU

### **PCT**

### **NOTIFICATION OF ELECTION**

(PCT Rule 61.2)

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Assistant Commissioner for Patents United States Patent and Trademark Office
Box PCT
Washington, D.C.20231
ETATS-UNIS D'AMERIQUE

Date of mailing (day/month/year)
23 August 2000 (23.08.00)

International application No.
PCT/EP99/10347

International filing date (day/month/year)
23 December 1999 (23.12.99)

In its capacity as elected Office

Applicant's or agent's file reference
PU9844

Priority date (day/month/year)
30 December 1998 (30.12.98)

ORLEFORS, Anna, Edman et al

Applicant

1.	The designated Office is hereby notified of its election made:
	X in the demand filed with the International Preliminary Examining Authority on:
	28 July 2000 (28.07.00)
	in a notice effecting later election filed with the International Bureau on:
2.	The election X was
i L	made before the expiration of 19 months from the priority date or, where Rule 32 applies, within the time limit under Rule 32.2(b).

The International Bureau of WIPO 34, chemin des Colombettes 1211 Geneva 20, Switzerland Authorized officer

Manu Berrod

Telephone No.: (41-22) 338.83.38

Facsimile No.: (41-22) 740.14.35

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According	y to international Patent Classification (IPC) or to both nation	and the same of	
B. FIELD	S SEARCHED		
Minimum TPC 7	documentation searched (classification system followed by C12Q B01L	classification symbols)	
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Daguera			
Document	ation searched other than minimum documentation to the e	xtent that such documents are included	in the fields searched
Electronic	data base consulted during the international search (name	of data base and, where practical, see	arch terms used)
İ			
	<u> </u>		
C. DOCUM	ENTS CONSIDERED TO BE RELEVANT		
Category •	Citation of document, with indication, where appropriate,	of the relevant passages	
			Relevant to claim N
A	WO 98 55653 A (NEXSTAR PHARM	IACEUTICALS	1-7
	INC  IO DECEMBER 1998 (1998-	12-10)	1-/
_ i	page 4 -page 5, line 22; cla		
A	WO 98 28440 A (DZIEGLEWSKA H	ANNA EVA	1 7
	I STIKUSEQUENCING AB (SF) NYP	EN PAAL (SE))	1-7
	2 July 1998 (1998-07-02) claims 1,6,7,15		
. 1			
A	WO 98 07019 A (KIEFFER HIGGI	NS STEPHEN G	1,2,8
Ī	;MIAN ALEC (US); KELLOGG GRE 19 February 1998 (1998-02-19	GORY (US); G)	1 -,-,0
	figures 11A-11E	,	
a	NO 07 47761 A (CARNOTE CONT.)		
`	WO 97 47761 A (SARNOFF CORP) 18 December 1997 (1997-12-18)		1-7
	page 11, paragraph 3; claims	/ 1~13	
ļ		-/	
V =			
	or documents are listed in the continuation of box C.	X Patent family member	ers are listed in annex.
	gories of cited documents:		
nemuoob "/ sebianco	t defining the general state of the art which is not red to be of particular relevance	or priority date and not in	after the international filing date conflict with the application but
earlier do	cument but published on or affective international	invention	inciple or theory underlying the
. document	Which may throw doubte an advance at a con-	"X" document of particular rele cannot be considered now	vance; the claimed invention vel or cannot be considered to
citation o	or other special reason (as specified)	"Y" document of particular mile	when the document is taken alone
		document is combined with	th one an inventive step when the
document later than	published prior to the international filing date but the priority date claimed	in the art.	neing abytique to a person skilled
	tual completion of the international search	"&" document member of the s	
		Date of malling of the inter	mational search report
	April 2000	26/04/2000	
me and mai	ling address of the ISA	Authorized officer	
	European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijewijk Tel. (+31-70) 340-2040, Tx. 31 651 epo ni,		
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C.(Continue	Rtion) DOCUMENTS CONSIDERED TO BE RELEVANT	PCT/EP 99	9/10347
Category *	Citation of document, with indication, where appropriate, of the relevant passages		Polovenia
,			Relevant to claim No.
,	WO 97 22825 A (NEUKERMANS ARMAND P) 26 June 1997 (1997-06-26) WO 98 45481 A (KNAPP MICHAEL ; BOUSSE LUC J		
	15 October 1998 (1998–10–15)		
	WO 97 21090 A (GAMERA BIOSCIENCE) 12 June 1997 (1997-06-12) cited in the application claims 46,57; figures 17A-E		1,2,8
		·	
	nitruation of ecoond cheet) (July 1922)		

# INTERNATIONAL SEARCH REPORT on patent family members

Inter. pplication No PCT/EP 99/10347

		1.		101721	33/1034/
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WO 9828440	A	02-07-1998	AU	5331198 A	17-07-1998
			EP.	. 0946752 A	06-10-1999
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			CN	1208464 A	17-02-1999
			EP	0865606 A	23-09-1998
•			NO	982563 A	05-08-1998
			AU	4144897 A	06-03-1998
			EP	0917648 A	26-05-1999
			WO	9807019 A	19-02-1998

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# **PCT**

REC'D 1 7 JAN 2001

# INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

Applicant's o	or agent's file reference		See Notification of Transmittal of International
PU9844		FOR FURTHER ACTION	Preliminary Examination Report (Form PCT/IPEA/416)
International	application No.	International filing date (day/montl	n/year) Priority date (day/month/year)
PCT/EP9	9/10347	23/12/1999	30/12/1998
International C12Q1/68		or national classification and IPC	
Applicant  AMERSH	AM PHARMACIA BIOT	FECH AB et al	
and is	transmitted to the applica	ant according to Article 36.	d by this International Preliminary Examining Authority
2. This R	EPORT consists of a total	al of 6 sheets, including this cover s	heet.
be (s	een amended and are the	basis for this report and/or sheets on 607 of the Administrative Instruct	ne description, claims and/or drawings which have containing rectifications made before this Authority ions under the PCT).
3. This re	eport contains indications	relating to the following items:	
l i	☑ Basis of the report		
11	☐ Priority		
111	☐ Non-establishment	of opinion with regard to novelty, in	ventive step and industrial applicability
IV	Lack of unity of inv		
٧	<ul><li>Reasoned stateme citations and expla</li></ul>	ent under Article 35(2) with regard to nations suporting such statement	novelty, inventive step or industrial applicability;
VI VI	☐ Certain document		
VII		the international application	
VIII	□ Certain observation	ns on the international application	
Date of sub	mission of the demand	Date o	f completion of this report
28/07/20	00	11.01.3	2001
Name and preliminary	mailing address of the international examining authority:	ational Author	zed officer
<u></u>	European Patent Office D-80298 Munich Tel. +49 89 2399 - 0 Tx: 5		n, A-C

Telephone No. +49 89 2399 8688

Fax: +49 89 2399 - 4465



International application No. PCT/EP99/10347

### I. Basis of the report

	the		on under Article 14 are referred to in this report as "originally filed" and are not annexed to o not contain amendments (Rules 70.16 and 70.17).):					
	1-16	6	as originally filed					
	Cla	ims, No.:						
	1-8		as originally filed					
	Dra	wings, sheets:						
	1/5-	5/5	as originally filed					
2.			guage, all the elements marked above were available or furnished to this Authority in the international application was filed, unless otherwise indicated under this item.					
	The	se elements were a	available or furnished to this Authority in the following language: , which is:					
		the language of a	translation furnished for the purposes of the international search (under Rule 23.1(b)).					
		the language of pu	ublication of the international application (under Rule 48.3(b)).					
		the language of a 55.2 and/or 55.3).	translation furnished for the purposes of international preliminary examination (under Rule					
3.			cleotide and/or amino acid sequence disclosed in the international application, the y examination was carried out on the basis of the sequence listing:					
		contained in the in	sternational application in written form.					
		filed together with	the international application in computer readable form.					
		furnished subsequ	ently to this Authority in written form.					
		furnished subsequently to this Authority in computer readable form.						
			It the subsequently furnished written sequence listing does not go beyond the disclosure in pplication as filed has been furnished.					
		The statement that listing has been full	It the information recorded in computer readable form is identical to the written sequence irnished.					
4.	The	amendments have	e resulted in the cancellation of:					
		the description,	pages:					
		the claims,	Nos.:					

1. This report has been drawn on the basis of (substitute sheets which have been furnished to the receiving Office in



International application No. PCT/EP99/10347

		the drawings,	sheets:		
5.					ome of) the amendments had not been made, since they have been as filed (Rule 70.2(c)):
		(Any replacement she report.)	eet contair	ning such	amendments must be referred to under item 1 and annexed to this
6.	Add	itional observations, if	necessar	y:	
٧.		soned statement und tions and explanation			rith regard to novelty, inventive step or industrial applicability; ch statement
1.	Stat	ement			
	Nov	relty (N)	Yes: No:		2,4,6,8 1,3,5,7
	Inve	entive step (IS)	Yes: No:	Claims Claims	- 1-8
	Indu	ustrial applicability (IA)	Yes: No:	Claims Claims	
2.	Cita	tions and explanations	3		

### VII. Certain defects in the international application

see separate sheet

The following defects in the form or contents of the international application have been noted: see separate sheet

### VIII. Certain observations on the international application

The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made: see separate sheet



### International application No. PCT/EP99/10347

### **EXAMINATION REPORT - SEPARATE SHEET**

### Re Item V

The following documents are referred to in this communication:

D1: WO 98 28440 A

cited in the application D2: WO 97 21090 A

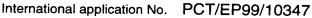
The present application is directed to a sequencing system which is based on a device comprising microchannels which are interconnected and wherein liquid flow is achieved by centripetal force.

#### 1 Novelty (Art 33(2) PCT):

#### Claim 1 is not novel over D1: 1.1

The method as defined in the preamble of the claim is known from D1 (D1: p 20 I 8-33) and the only characterising feature of the invention is the "microfluidic device". This expression however is a designation which does not contain the essential technical features of the invention. A "microfluidic device" can mean any device which is suitable to operate with liquids on a microlitre scale. Thus, a microtitre plate for example as disclosed in D1 (p 13 para 3 - p 14 para 3) falls within the meaning of "microfluidic device". Since D1 also discloses the detection of PP release by light emission from a luciferin luciferase reaction (D1: p 7 para 3 ff), claim 5 is not novel, either. Similarly, D1 also discloses real-time monitoring of the sequencing reaction (D1: p 3 para 2; p 8 last paragraph - p 10 para 1) and therefore anticipates the subject-matter of claim 7.

- 1.2 The subject-matter of claim 3 is not novel, either. All the method steps are anticipated by D1 (D1: p 20 I 8-33). The range of immobilized single stranded DNA (step (i)) covers the usual range of primers used for an enzymatic sequencing reaction. The same applies to the range given in step (ii). The "predetermined areas on the surface of a microfluidic device" is not a distinguishing feature over D1 either, because wells of a microtitre plate as described in D1 (D1: p 13 para 4) fall within the meaning of said expression.
- 1.3 Claim 2 is novel, because none of the available documents describes a



**EXAMINATION REPORT - SEPARATE SHEET** 

sequencing method in which the sample DNA is in a first step immobilized on a reaction area in a microchannel structure of a microfluidic device and after that a deoxynucleotide is added before the template dependent incorporation of the deoxynucleotide is detected.

For the same reason, also claims 4, 6 and 8 are novel.

#### 2 Inventive Step (Art 33(3) PCT):

Claim 2 does not appear to be inventive for the following reasons: 2.1

D1, which is considered to represent the closest prior art, describes a method of sequencing DNA based on the detection of the release of pyrophosphate. An embodiment is disclosed (D1: p 20 I 8-33) which shows all the method steps but is not carried out in reaction areas of a microchannel structure.

This distinguishing feature however, does not appear to solve a technical problem. Reaction areas in a microchannel structure seem to be the equivalent of e.g. wells on a microtitre plate. Since the claimed method does not rely on the microchannel structures, it does not achieve an effect over the method of D1.

Thus, claim 2 does not seem to satisfy Art 33(3) PCT.

2.2 Claim 4 does not appear to be inventive for the following reasons:

D1, which is considered to represent the closest prior art, is distinguished from the subject-matter of claim 4 in that D1 does not disclose a method in which the sample DNA is moved through a microfluidic device before being immobilized in the reaction chamber. This distinguishing feature does not seem to solve a problem. Since dependent claims 6 and 8 do not contain an inventive concept per se, they do not appear to comply with Art 33(3) PCT, either.

Claim 8 contains additional features of the microfluidic device. However, the use of centripetal forces to motivate fluid movement through microchannels are known from D2 (abstract). The disclosure of D2 also embraces the use of such devices for DNA sequencing (D2: Example 7) and various other methods for DNA analysis (D2: Examples 3-6).

### Re Item VII

In order for the application to be self-contained the reference to non-published patent



**EXAMINATION REPORT - SEPARATE SHEET** 

applications should have been replaced by the corresponding publication numbers (Guidelines II 4.17)(e.g. p 5 I 26; p 6 I 1).

### Re Item VIII

- 1 The claims lack essential features of the invention (Art 6 PCT): In the description it is stated that the advantage of the method lies in the small amounts of reagents needed (application: p 10 | 29-31). However, none of the present claims contains a technical feature defining the volumes used in the "microfluidic device".
- 2 The use of brackets in claims 1 and 2 renders the scope of said claims unclear (Art 6 PCT) since it is not clear if the features in the brackets are meant to belong to the claims. Moreover, the question mark in claim 1(i) renders the claim unclear.
- 3 Claim 2 does not comply with Art 6 PCT for the following reason: Step (i) relates to "forming immobilised double stranded DNA [...] and step (ii) specifies that the "double stranded DNA" consists of a sample DNA strand and a primer strand. This does technically not make sense, because a primer should be shorter than the sample DNA strand in order to enable the template dependent primer extension, which means that the DNA molecule can only be partly double stranded. The DNA molecule must have a single stranded overhang.
- 4 Claim 3 does not satisfy Art 6 PCT, because its subject-matter is not clearly defined: According to the preamble the claim is directed to a method of determining a nucleotide base in a sample whereas step (v) relates to determining the sequence of portions of sample DNA. Moreover, the dependency of the claim is unclear: According to the preamble of the claim, it is directed to a method of determining a nucleotide base whereas claim 2 is directed to the identification of a sequence of a portion of sample DNA.
- 5 Claim 4 is not clear (Art 6 PCT): Step (v) relates to "extending the primer [...] with a known deoxynucleotide [...] or dideoxynucleotide [...]". This formulation implies that the person who carries out the method already knows which nucleotide to take. This, however would render the whole method pointless since it is aimed at identifying the sequence.

 $\lambda \geqslant 1$ 

# 001 08:37 \* Pg 17/22 NR.689

# **PCT**

**PATENT COOPERATION TREATY** 

## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

Applicant's	жаде	nt's file reference	FOR FURTHER ACTION		ation of Transmittal of International
PU9844			FOR FORTHER ACTION	Premmary	Examination Report (Form PCT/IPEA/416)
Internationa	appli	cation No.	International filing date (day/mont	h/year)	Priority date (day/month/year)
PCT/EP9	9/10:	347	23/12/1999		30/12/1998
Internationa C12Q1/66		nt Classification (IPC) or I	national classification and IPC		
Applicant AMERSH	AM I	PHARMACIA BIOTE	CH AB et al		
1. This ir and is	terna trans	ational preliminary examitted to the applicant	mination report has been prepare according to Article 36.	d by this Inte	emational Preliminary Examining Authority
2. This P	EPO	RT consists of a total	of 6 sheets, including this covers	sheet.	
be	en a	mended and are the b	ied by ANNEXES, i.e. sheets of t asis for this report and/or sheets 607 of the Administrative Instruct	containing re	on, claims and/or drawings which have actifications made before this Authority ne PCT).
These	ann	exes consist of a total	of sheets.		
3. This r			elating to the following items:		
1	<b>23</b>	•			
11					No. 4 The
311			opinion with regard to novelty, in	iventive step	and industrial applicability
17				novothe inv	entive step or industrial applicability;
٧	_		tions suporting such statement	Tioveny, into	entive step of modelital applicability,
VI		Certain documents	•		
VII			international application		
VIII	Ø	Certain observations	on the international application		
Date of sub	missi	on of the demand	Date o	f completion o	f this report
28/07/20	00		11.01.	2001	
	exam	g address of the internation	onal Author	tzed officer	S TO S MONTH OF THE PARTY OF TH
<u> </u>	D-8	opean Patent Office 0298 Munich +49 89 2399 - 0 Tx: 523		m, A-C	
		: +49 89 2399 - 4465	· · · · · · · · · · · · · · · · · · ·	one No. +49 8	39 2399 8688

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		is of the report						
1.	resp the i	This report has been drawn on the basis of (substitute sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to the report since they do not contain amendments (Rules 70.16 and 70.17).): Description, pages:						
	1-16	as originally filed						
	Clai	ms, No.:						
	1-8	as originally filed						
	Dra	wings, sheets:						
	1/5-	5/5 as originally filed						
2.	lang	n regard to the <b>language</b> , all the elements marked above were available or furnished to this Authority in the guage in which the international application was filed, unless otherwise indicated under this item.						
	The	se elements were available or furnished to this Authority in the following language: , which is:						
		the language of a translation furnished for the purposes of the international search (under Rule 23.1(b)). the language of publication of the international application (under Rule 48.3(b)).						
		the language of a translation furnished for the purposes of international preliminary examination (under Rule 55.2 and/or 55.3).						
3.	Witl inte	n regard to any nucleotide and/or amino acid sequence disclosed in the international application, the mational preliminary examination was carried out on the basis of the sequence listing:						
		contained in the international application in written form.						
		filed together with the international application in computer readable form.						
		furnished subsequently to this Authority in written form.						
		furnished subsequently to this Authority in computer readable form.						
		The statement that the subsequently furnished written sequence listing does not go beyond the disclosure in the international application as filed has been furnished.						
		The statement that the information recorded in computer readable form is identical to the written sequence listing has been furnished.						
4	The	amendments have resulted in the cancellation of:						

the description,

pages:

Nos.:

NR.689

	the drawings,	sheets:
5.	This report has been considered to go bey	established as if (some of) the amendments had not been made, since they have been nond the disclosure as filed (Rule 70.2(c)):
	(Any replacement sh report.)	eet containing such amendments must be referred to under item 1 and annexed to this

- 6. Additional observations, if necessary:
- V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement
- 1. Statement

. .

Yes: Claims 2,4,6,8 Novelty (N) Claims 1,3,5,7 No: Yes: Claims -Inventive step (IS) No: Claims 1-8 Claims 1-8 Yes: Industrial applicability (IA)

No:

2. Citations and explanations see separate sheet

### VII. Certain defects in the international application

The following defects in the form or contents of the international application have been noted: see separate sheet

Claims -

### VIII. Certain observations on the international application

The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made: see separate sheet

P020

**EXAMINATION REPORT - SEPARATE SHEET** 

### Re Item V

The following documents are referred to in this communication:

D1: WO 98 28440 A

D2: WO 97 21090 A cited in the application

The present application is directed to a sequencing system which is based on a device comprising microchannels which are interconnected and wherein liquid flow is achieved by centripetal force.

#### 1 Novelty (Art 33(2) PCT):

#### 1.1 Claim 1 is not novel over D1:

The method as defined in the preamble of the claim is known from D1 (D1: p 201 8-33) and the only characterising feature of the invention is the "microfluidic device". This expression however is a designation which does not contain the essential technical features of the invention. A "microfluidic device" can mean any device which is suitable to operate with liquids on a microlitre scale. Thus, a microtitre plate for example as disclosed in D1 (p 13 para 3 - p 14 para 3) falls within the meaning of "microfluidic device". Since D1 also discloses the detection of PP, release by light emission from a luciferin luciferase reaction (D1: p 7 para 3 ff), claim 5 is not novel, either. Similarly, D1 also discloses real-time monitoring of the sequencing reaction (D1: p 3 para 2; p 8 last paragraph - p 10 para 1) and therefore anticipates the subject-matter of claim 7.

- 1.2 The subject-matter of claim 3 is not novel, either. All the method steps are anticipated by D1 (D1: p 20 I 8-33). The range of immobilized single stranded DNA (step (i)) covers the usual range of primers used for an enzymatic sequencing reaction. The same applies to the range given in step (ii). The "predetermined areas on the surface of a microfluidic device" is not a distinguishing feature over D1 either, because wells of a microtitre plate as described in D1 (D1: p 13 para 4) fall within the meaning of said expression.
- 1.3 Claim 2 is novel, because none of the available documents describes a

**EXAMINATION REPORT - SEPARATE SHEET** 

sequencing method in which the sample DNA is in a first step immobilized on a reaction area in a microchannel structure of a microfluidic device and after that a deoxynucleotide is added before the template dependent incorporation of the deoxynucleotide is detected.

For the same reason, also claims 4, 6 and 8 are novel.

- 2 Inventive Step (Art 33(3) PCT):
- 2.1 Claim 2 does not appear to be inventive for the following reasons:

D1, which is considered to represent the closest prior art, describes a method of sequencing DNA based on the detection of the release of pyrophosphate. An embodiment is disclosed (D1: p 20 l 8-33) which shows all the method steps but is not carried out in reaction areas of a microchannel structure.

This distinguishing feature however, does not appear to solve a technical problem. Reaction areas in a microchannel structure seem to be the equivalent of e.g. wells on a microtitre plate. Since the claimed method does not rely on the microchannel structures, it does not achieve an effect over the method of D1.

Thus, claim 2 does not seem to satisfy Art 33(3) PCT.

2.2 Claim 4 does not appear to be inventive for the following reasons:

D1, which is considered to represent the closest prior art, is distinguished from the subject-matter of claim 4 in that D1 does not disclose a method in which the sample DNA is moved through a microfluidic device before being immobilized in the reaction chamber. This distinguishing feature does not seem to solve a problem. Since dependent claims 6 and 8 do not contain an inventive concept per se, they do not appear to comply with Art 33(3) PCT, either.

Claim 8 contains additional features of the microfluidic device. However, the use of centripetal forces to motivate fluid movement through microchannels are known from D2 (abstract). The disclosure of D2 also embraces the use of such devices for DNA sequencing (D2: Example 7) and various other methods for DNA analysis (D2: Examples 3-6).

### Re Item Vil

In order for the application to be self-contained the reference to non-published patent

**EXAMINATION REPORT - SEPARATE SHEET** 

applications should have been replaced by the corresponding publication numbers (Guidelines II 4.17)(e.g. p 5 l 26; p 6 l 1).

### Re Item VIII

- The claims lack essential features of the invention (Art 6 PCT): In the description it 1 is stated that the advantage of the method lies in the small amounts of reagents needed (application: p 10 I 29-31). However, none of the present claims contains a technical feature defining the volumes used in the "microfluidic device".
- The use of brackets in claims 1 and 2 renders the scope of said claims unclear 2 (Art 6 PCT) since it is not clear if the features in the brackets are meant to belong to the claims. Moreover, the question mark in claim 1(i) renders the claim unclear.
- Claim 2 does not comply with Art 6 PCT for the following reason: Step (i) relates to "forming immobilised double stranded DNA [...] and step (ii) specifies that the "double stranded DNA" consists of a sample DNA strand and a primer strand. This does technically not make sense, because a primer should be shorter than the sample DNA strand in order to enable the template dependent primer extension, which means that the DNA molecule can only be partly double stranded. The DNA molecule must have a single stranded overhang.
- Claim 3 does not satisfy Art 6 PCT, because its subject-matter is not clearly defined: According to the preamble the claim is directed to a method of determining a nucleotide base in a sample whereas step (v) relates to determining the sequence of portions of sample DNA. Moreover, the dependency of the claim is unclear: According to the preamble of the claim, it is directed to a method of determining a nucleotide base whereas claim 2 is directed to the identification of a sequence of a portion of sample DNA.
- Claim 4 is not clear (Art 6 PCT): Step (v) relates to "extending the primer [...] with 5 a known deoxynucleotide [...] or dideoxynucleotide [...]". This formulation implies that the person who carries out the method already knows which nucleotide to take. This, however would render the whole method pointless since it is aimed at identifying the sequence.

### **PCT**





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(54) Title: METHOD FOR SEQUENCING DNA USING A MICROFLUIDIC DEVICE

### (57) Abstract

The present invention describes a method for sequencing DNA in a device with microfluidics properties and a set of reagents for its use. This microfluidic device may be in form of a disc with radially extending microchannel structures (CD form) having an inner application area that may be common for one or more microchannel structures. By spinning the disc the liquid can be driven from an applicator area into reaction and/or detection are as closer to the periphery of the disc. Liquid transportation may thus be driven by centripetal force. The microfluidic device may also have other gemoetrical forms. Several methods can be used to determine the sequence of DNA according to the invention but the real time determination of released pyrophosphate using the luciferase luciferin reaction is preferred.

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* Special car	tegories of cited documents :	"T" later document published	after the international filing date		
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WO 00/40750



09/869554 Recyclypto 2 8 JUN 2001

### METHOD FOR SEQUENCING DNA USING A MICROFLUIDIC DEVICE

A method, device and reagents for the high throughput sequencing of nucleic acids.

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This invention is based on a method for sequencing nucleic acids, a device for handling DNA containing samples, and a reagent kit, where the sequencing method is based on pyrosequencing.

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DNA sequencing is an essential tool in basic molecular biology research. In the future it can be expected that DNA sequencing will be used in both diagnostic research as well as applied genome diagnostics.

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The majority of *de novo* DNA sequencing is carried out with shot gun sequencing and with the enzymatic chain terminating method of Sanger. The sequence is generated by the resolution, using gel electrophoresis, of DNA fragments which have been prepared by elongating predetermined oligonucleotide primers. The separation of DNA fragments and the following analysis are cumbersome and great efforts have been made to automate these steps. Despite the fact that automated DNA sequencers are used in large scale genome projects there is a need for DNA sequencing devices with higher throughput, for both genome sequencing and routine clinical applications.

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Pyrosequencing is a modified pyrophosphate (PPi)-based sequencing method in which PPi is detected by the release of light in the luciferase – luciferin reaction (see for example PCT patent applications WO 98/13523 and 98/28440). Each time one nucleotide molecule is incorporated into the growing DNA strand one molecule of PPi is released. The light detected is directly proportional to the number of incorporated bases in the growing DNA strand. The main drawback with this method is the number of samples that can be handled simultaneously and the speed of detection. Thus PCT application WO 98/28440 describes reactions in 96 well microtitre plates. Since the volume in each well is between 10 - 500 microlitre, the costs for the reagents are high and limit the use of the method.

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When performing the pyrosequencing method in microtitre plates reaction mixes are added to the reaction chamber, but since no solution is removed from the well, the reaction can only be done a limited number of times, thereby only generating short stretches of DNA sequences. One of the major problems is to remove the excess of dNTP that can lead to misincorporation and dATP, which interferes with the light generation reaction. WO 98/28440 describes the addition of a nucleotide degrading enzyme, e.g. apyrase, to deal with this.

It is an object of the present invention to overcome several of the previous problems with pyrosequencing, such as the increasing volume when performing pyrosequencing in a microtitreplate, as well as reducing the consumption of reagents and making it feasible for analysis of several hundred samples simultaneously, thus providing a high through-put system.

Arrayed Primer Extension (APEX), works by immobilising a large number of primers to a solid surface, thus creating a DNA-chip. These primers are constructed to be consecutively overlapping over the entire gene of interest, so that every base in the gene will have a primer to its 5'-end. By adding fluorescently labelled dideoxynucleotides, the primers will then be extended by one nucleotide using the sample DNA as template. It will thus be easy to check which nucleotide was incorporated, which in turn tells you the entire sequence of the sample DNA.

The present invention describes a method for sequencing DNA in a device with microfluidics properties and a set of reagents for its use. This microfluidic device may be in form of a disc with radially extending microchannel structures (CD form) having an inner application area that may be common for one or more microchannel structures. By spinning the disc the liquid can be driven from an applicator area into reaction and/or detection are as closer to the periphery of the disc. Liquid transportation may thus be driven by centripetal force. The microfluidic device may also have other geometrical forms.

Accordingly, in a first aspect the present invention provides a method for identifying the sequence of a portion of sample DNA, which method comprises:

- forming immobilised double stranded DNA on one or more reaction areas
   in a microchannel structure of a microfluidic device;
- (ii) adding a known deoxynucleotide, (or the corresponding deoxynucleotide analogue or dideoxynucleotide) and a DNA polymerase to each of said one or more reaction areas so that extension of primer only occurs if there is a complementarity of the added deoxynucleotide or dideoxynucleotide with the strand of sample DNA that is part of the immobilised double stranded DNA;
- (iii) detecting whether or not the deoxynucleotide, deoxynucleotide analogue or dideoxynucleotide added in step (ii) has been added to the primer DNA in said one or more reaction areas,
- (iv) repeating steps (ii) and (iii) as required with a different deoxynucleotide (or the corresponding deoxynucleotide analogue or dideoxynucleotide).

The double stranded DNA which after step (i) is present in the reaction area consists of one strand of primer DNA and one strand of sample DNA (template). One of the strands is firmly attached to the reaction area. At least one of the strands of sample DNA and primer DNA is different for at least two reaction areas within one and the same microfluidic device.

The immobilised double stranded DNA comprising template and primer may be formed outside the microfluidic device as described in the experimental part. In the most efficient variants, it is, however, believed that the immobilised double strand is formed within the microfluidic device, for instance in the reaction chamber, by introducing separately either single or double stranded sample DNA and primer DNA. In case double stranded sample DNA is introduced in step (i) above, or in the preferred aspects described later, it has to be denatured within the microfluidic device.

The added deoxynucleotide, deoxynucleotide analogue or dideoxynucleotide in step (ii) may be labelled or unlabelled. If it is labelled the label as such is measured. Any kind of label that can be incorporated in a nucleic acid strand by the polymerase can be used, for instance a fluorescent label. If the

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deoxynucleotide, deoxynucleotide analogue or dideoxynucleotide is unlabelled, nucleotide incorporation can be detected by measuring the amount of PP<sub>i</sub> released.

The amount of single stranded sample DNA that is immobilised is typically 0.1-200 pmole but may also be as low as 1 atomole, for example 1 femtomole. The number of reaction areas may be from two upwards. Typically it is below 500,000 such as below 100,000.

In both the general and preferred aspects, the length of the elongated part of the primer may be from one base upwards. In case the method is arrayed primer extension (APEX), WO 95/00699, the elongated part of the primer is one nucleotide, for example, when using a labelled terminator e.g. dideoxynucleotide. This means that the repeating step (iv) is run at most three times.

In one aspect the present invention comprises:

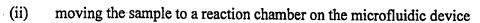
- attaching 0.1 200 pmol of a primer or single stranded DNA sample to each of between one and 100,000 pre-determined areas on the surface of a microfluidic device;
- (ii) hybridising small amounts, e.g. 0.1 200 pmol, of single stranded sample DNA or primer respectively to each of the predetermined areas;
- (iii) adding a known deoxynucleotide, deoxynucleotide analogue or dideoxynucleotide and a DNA polymerase so that extension of the primer only occurs, with consequent release of pyrophosphate (PPi), if there is a complementarity with the sample DNA;
- (iv) measuring the release of PPi and from which predetermined area on the device it is released;
- (v) repeating steps (iii) and (iv) as required to construct a DNA sequence for the elongated primers, and hence for portions of the sample DNA.

In a preferred aspect, the present invention provides a method for identifying the sequence of a portion of sample DNA, which method comprises:

(i) adding sample DNA to a predetermined area on a microfluidic device

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- (iii) attaching the sample DNA to a surface of the reaction chamber, alternatively hybridising the sample DNA in a single stranded form to a primer attached to the reaction chamber (then to (v))
- (iv) if the sample DNA has not been attached to a primer attached to the reaction chamber, hybridising a primer to the DNA in a single stranded form
- (v) extending the primer in the presence of a DNA polymerase with a known deoxynucleotide (dNTP), deoxynucleotide analogue or dideoxynucleotide (ddNTP), such extension being indicated by detection of pyrophosphate (PPi) released from the extension reaction
- (vi) repeating step (v) as required to establish the sequence of the extended primer.
- The sample DNA to be loaded onto the microfluidic device may be an amplified sample and/or may be amplified within the microfluidic device. Amplification may involve introduction of a tag suitable for attaching the amplified DNA to a solid support.
- The present invention may be applied to all fields where DNA or RNA are sequenced. These are *de novo* sequencing, resequencing of known sequences for monitoring mutation or base changes, sequencing of sequence polymorphisms and mini-sequencing where only one base is determined (including arrayed primer extension (APEX). Furthermore, the present invention may be applied to situations where the identity of a number of polymorphs is determined at the same time (see for example European Patent application 99303215.0).

The DNA to be sequenced can be of any origin: animal, plant, bacterial, or viral.

This DNA can be amplified either in the device or before it is loaded onto the device.

The microfluidic device of the present invention may be analogous to those described in the literature, see patent application WO97/21090 filed by Gamera BioScience, and is preferably in the form of a disc, where the fluids are moved by

### WO 00/40750

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centripetal forces see for example co-pending application GB 9809943.5. The device preferably has a sample loading or application area with one or more reaction chambers and a detection chamber. Thus, a reaction can be performed in the detection chamber, and any light reaction can be detected directly when it occurs. In the case of separated chambers the flow between these can be steered by different types of barriers, like narrowed transport channels, different mechanical barriers or by surface interactions between the walls and the solution. These interactions can be of hydrophobic – hydrophilic character.

### 10 Kit of reagents:

Buffers for amplification

(vii) Enzymes for amplification, or mixed with the buffer Sequencing buffer, luciferin Sequencing enzymes, can also be mixed with the buffer dCTP, dGTP, dTTP in separate buffers dATPαS in buffer

These reagents can also be stored in a dried state e.g. glacified, direct in the disc and the reagents will then be activated first after addition of water.

### Enzymes to be used in the kit:

DNA polymerase or another thermostable DNA polymerase for amplification and/or sequencing reactions e.g. Taq or other thermostable DNA polymerases ATP sulphurylase

25 Luciferase

Apyrase as an optional non-preferred ingredient

Illustrative DNA polymerases are Klenow fragment polymerases. Sequenases and other 3'-5' exo- DNA polymerases, and Taq DNA polymerases and other thermostable polymerases. 3'-5' exo- DNA polymerases are preferred.

Amplification reactions on sample DNA may be performed within the microfluidic device or outside it before the sample DNA is loaded onto the device.

A kit according to the invention comprises a microfabricated device, preferably in

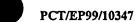
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form of a disc with radially extending microchannel structures, in combination with one, two or three of (a) Luciferase, (b) DNA polymerase, and (c) ATP sulfurylase, optionally combined with any of the above-mentioned ingredients, with preference for one or more ingredients that relate to anyone of (a)-(b) as a substrate.

The method of loading reagents and liquids to the microfluidic device can be with a dispenser, or a mechanical device for "picking" the different samples. The loading device should be able to load the different application spots onto the spinning device, preferably during the spinning of the device.

After the loading of the sample, it will be transferred by the centripetal force to the reaction chamber. In the reaction chamber the sample should be attached to a wall of the reaction chamber. If the sample is a DNA fragment, it can be attached to a surface of the reaction chamber in one of the following ways.

In the first, the DNA will be tagged at the 3'- or 5'- end during the amplification step, the tag can preferably be biotin or any other suitable tag described in the literature and suitable for attaching the tagged substance to a solid support. The surface in the reaction chamber should be activated with a substance to quickly and effectively bind to the DNA tag, preferably streptavidin will be used when the tag is biotin. The surface in the chamber can also be enlarged by the use of beads or other surface enlargement groups or structures, for example agarose or polystyrene-divinyl benzene beads (Sepharose or Source, respectively, Amersham Pharmacia Biotech AB) that are retained in the chamber, for instance by being glued to the wall of the reaction chamber. The beads or the enlargement groups may then carry the appropriate affinity group for catching the tagged DNA, for instance strepavidin in case the tag is biotin.

A second way to bind the DNA to the surface is by attaching the primer before amplification and then to perform the amplification of the sample DNA in the reaction chamber. With this approach additional coupling chemistries can be used to link the primer to the surface, such as an aminolinker on the primer with an epoxysilane treated surface.

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A third way is to select the DNA sample of interest with attached primer(s). These can then be used for both attachment and act as sequencing primers, simultaneously. With this approach, the sample DNA needs to be fragmented prior to the hybridisation to the primers in the reaction chamber, following this hybridisation, the primer is extended. An advantage with this method is that several or many hundred different primers can be attached in the reaction chamber and these can be made in such a way that they cover different parts of the DNA fragment of interest, thereby the whole DNA fragment can be sequenced in one step. The distance on an unfragmented DNA molecule between sequences binding to different primers can vary between 1 to 500 bases and is most preferably 5-50 bases apart.

Binding of DNA to the reaction area may be by covalently linking one of the strands, preferably the primer, directly to the surface of a reaction area or via a specific adsorption such as via biotin-avidine as described above and other affinity pairs providing a sufficient binding to each other. A number of techniques for covalently linking DNA to solid supports are known in the scientific and patent literature.

When the sample DNA is attached to the surface it should be denatured, this can be achieved by several methods for example, hydrogen bond breaking agents, high pH or high temperature. In the present invention the preferred method is denaturing the DNA with high pH, preferably by using sodium hydroxide. Denaturation can take place either outside or inside the microfluidic device.

The following step in the invention is the elongation where DNA polymerase is added together with primer; optionally the primer can be added prior to the other compounds. The other reagents are ATP sulphurylase, luciferase, L- and D-luciferin and APS and one of the nucleotide triphosphates, dATP $\alpha$ S, dCTP, dGTP or dTTP. These are added sequentially, i.e. a mix with dATP $\alpha$ S and the other reagents, followed by a detection step and finally a wash, this is followed by dC, and then dG, and then dT or any other predetermined order. When a nucleotide is incorporated a signal is detected in the luciferase reaction and this is scored as that



base. The washing step included here solves the problem with loading the reaction mixes to one well many times and thereby getting a larger and larger volume. Since the washing here is included in the spinning device there is no need for the use of apyrase as described in Patent application WO 98/28440.

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PPi can be determined by many different methods and a number of enzymatic methods have been described in the literature (Reeves *et al.*, (1969), Anal. Biochem., 28, 282-287; Guillory *et al.*, (1971), Anal. Biochem., 39, 170-180; Johnson *et al.*, (1968), Anal. Biochem., 15, 273; Cook *et al.*, (1978), Anal. Biochem., 91, 557-565; and Drake *et al.*, (1979), Anal. Biochem., 94, 117-120).

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It is preferred to use luciferase and luciferin in combination to quantify the release of pyrophosphate since the amount of light generated is substantially proportional to the amount of pyrophosphate released which, in turn, is directly proportional to the amount of base incorporated. The amount of light can readily be estimated by a suitable light sensitive device such as a luminometer, or a photomultiplying device in close proximity to the device of the present invention.

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Luciferin-luciferase reactions to detect the release of PPi are well known in the art. In particular, a method for continuous monitoring of PPi release based on the enzymes ATP sulphurylase and luciferase has been developed by Nyrén and Lundin (Anal. Biochem., 151, 504-509, 1985) and termed ELIDA (enzymatic Luminometric Inorganic Pyrophosphate Detection Assay). The method may however be modified, for example by the use of a more thermostable luciferase (Kaliyama *et al.*, 1994, Biosci. Biotech. Biochem., 58, 1170-1171) and/or ATP sulfurylase (Onda *et al.*, 1996, Bioscience, Biotechnology and Biochemistry, 60:10, 1740-42). This method is based on the following reactions:

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ATP sulphurylase PPi + APS  $\longrightarrow$  ATP + SO<sup>2</sup>-4

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### luciferase

ATP + luciferin + 
$$O_2$$
 -----> AMP + PPi + oxyluciferin +  $CO_2$  +  $hv$ 

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### (APS = adenosine 5'-phosphosulphate)

The preferred detection enzymes involved in the PPi detection reaction are thus ATP sulphurylase and luciferase.

The method of the invention may be performed in two steps, as described for example in WO 93/23564 and WO 89/09283, firstly a polymerase reaction step, i.e. a primer extension step, wherein the nucleotide(s) are incorporated, followed by a second detection step, wherein the release of PPi is monitored or detected, to detect whether or not a nucleotide incorporation has taken place. Thus, after the polymerase reaction has taken place, samples from the polymerase reaction mix may be removed and analysed by the ELIDA e.g. by adding an aliquot of the sample to a reaction mixture containing ELIDA enzymes and reactants.

However, as mentioned above, the method of the invention may readily be modified to enable the sequencing (i.e. base incorporation) reactions to be continuously monitored in real time. This may simply be achieved by performing the chain extension and detection, or signal-generation, reactions substantially simultaneously by including the "detection enzymes" in the chain extension reaction mixture.

The reaction mix for the polymerase reaction may thus include at least nucleotide (deoxy- or dideoxy), polymerase, luciferin, APS, ATP sulphurylase and luciferase together with an optional nucleotide-degrading enzyme e.g. apyrase. The polymerase reaction may be initiated by addition of the polymerase or, more preferably the nucleotide. Preferably the detection enzymes are already present at the time the reaction is initiated, or they may be added with the reagent that initiates the reaction.

With the use of a microfluidic system the volumes of reagents are in the range of nanolitres compared to microlitres in the 96 well format. This will reduce the consumption of reagents a thousand fold or more.

The present invention is illustrated by the following figures, which are by way of example only, wherein:

#### WO 00/40750

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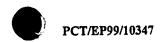


Figure 1: A schematic drawing of the fluidic channels in the spinning device. Samples are loaded either by a mechanical device or a piezo dispenser. The reaction chamber and the detection chamber can be the same. The outlet from the chamber/s will have some barrier to stop the fluid to move during the spinning. This barrier can be a hydrophobic surface.

Figure 2: Loading a CD from a liquid train. The train is loaded from the MTP (microtiter plate) and each sample (black in the diagram) is separated with air or an inert solution, a wash solution can also be included. When the whole plate is loaded in the capillary tube, a high pressure is applied in the same or opposite flow direction and samples will then be dispensed through the piezo dispenser on to application areas of the CD surface.

Figure 3: A schematic drawing of a mechanical device for loading a spinning device. The microtitre plate at the left, a wash station in the middle and the CD where samples, reagents and liquids should be applied to. 1) start position and transfer to the CD, 2) wash the applicator means, and 3) pick up new samples. The applicator means may be in the form of pens or syringes.

Figure 4a-d: show various parts and enlargements of the microchannel structures that have been used for proof of the principle utilised in the present invention.

Figure 4a: shows the peripheral part of a circular disc. The shown part have five microchannel structure extending radially outwards.

Figure 4b: shows an enlarged view of microchannel structure K9.

Figure 4c: shows an enlarged view of the sample volume definition unit in a microchannel structure.

Figure 4d: shows an enlarged view of the reaction chamber area plus chambers for disposal of waste liquids. In particular this figure indicates variations in depth (shadowed parts I, II, III and IV).



#### 1) Materials/Investigated units

5 Polymerase: Klenow Fragment (3' -5'exo-) New England Biolabs (storing buffer:

10mM Tris (pH 7,4) 1mM EDTA, 1mM DTT, 50% glycerol) 5U/μl or 50U/μl:

Pyrosequencing AB

Luciferase: Promega (13,33mg/ml)

Sulphurylase: Sigma (50mU/µl)

10 PolyvinylpyrollidonePVP: Sigma

MgAc<sub>2</sub>: Merck

D-Luciferin: BioThema

DTT: Sigma

Adenosines 5' phosphosulphate (APS): Sigma

15 dATPαS: Amersham Pharmacia Biotech

PPase pyrophosphatase: Sigma

dCTP, dTTP, dGTP: Amersham Pharmacia Biotech (PPi free)

Working solutions:

20 10 x stockA: PVP (4mg/ml); MgAc2 (10.7mg/ml); D-luciferin (1.0μg/ml); DTT

(1.0mM); Tris-Ac pH7.6(0.01M) APS (10mM)

Nucleotides: dATPaS(1.25mM), dCTP(0.5mM), dGTP(0.5mM), dTTP(1.25mM)

Buffers: Binding Washing Buffer (BW): 1M NaCI, 5mM Tris-HCI (pH 7.5),

0.5mM EDTA

TE Buffer: 10mM Tris-HCI, 1mM EDTA (pH 7.6)

TAE Buffer: 0,04M Tris-Acetate (7.8), mM EDTA

Particles: Source beads to which strepavidin has been attached; 15µm (APB)

Streptavidin (SA): 10mg/ml

Templates / Oligos

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Template: a 50-mer oligonucleotide with known sequence tagged with biotin in the 3' end.

Primer: 24-mer oligonucleotide complementary to part of the template.

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#### Microfluidic device material and treatment:

The microchannel structures (K7-K12) in figures 4a-d are arranged radially on a microfluidic disc. They start from a common annular inner application channel (1) and end in a common annular outer waste channel (2), coaxial with channel (1). Each inlet opening (3) of the microchannel structures may be used as an application area. Each microchannel structure is provided with a waste chamber (4) that opens into the outer waste channel (2). The flow direction is from the inlet openings (1) to the waste chamber (4). Flow is driven both by capillary action and centripetal force, i.e. by spinning the disc. Radial waste channels (5) directly connecting the annular inner channel (1) with the annular outer waste channel (2) are also shown.

Liquid passes from the inlet opening (3) via an entrance port (6) into a volume defining unit (7) and from there to a reaction chamber (10). The volume defining unit (7) has a passage into a waste channel (8) for removing excess liquid, e.g. to the annular outer waste channel (2), and a vent (9) which opens into open air. The reaction chamber (10) may become shallower (I,II,III,IV) (Fig 4d and Table) at the outlet end. A restricted channel (11) is provided between the reaction chamber (10 and the waste chamber (4). Due to the relatively large width of the waste chamber (4), there are preferably one or more supports (12) to ensure the rigidity of the chamber.

The volume defining unit (7) is U-shaped as shown in figure 4a-c with the entrance port (6) opening into the top of one of the legs of the U and the waste channel (8) starting from the other leg of the U, with a vent (9) placed at the top of this other leg. The bottom of the U-formed volume defining unit (7) is connected to the reaction chamber (10).

In addition to the application area at the inlet (3) of the structure, there may also be an additional application area (13) connected to the entrance port (6).

There is preferably also a vent (14) to open air in the reaction chamber (10). A

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hydrophobic break is preferably provided at the connection (16) of the reaction chamber (10) to the volume defining unit (7).

The outer annular waste channel (2) may be sectioned so as to collect waste from a predetermined number of closely located microchannel structures.

Hydrophobic breaks were introduced by marking with an over-head pen (permanent ink) (Snowman pen, Japan): (a) between microchannel structure inlets (3) in the inner annular application channel (1), (b) each opening (15) into the outer annular waste channel (i.e. the openings of the waste chambers) and, (c) if present, also the radial waste channels (5) which connect the inner annular application channel (1) and the outer annular waste channel (2), and also the waste channel (8) which guides away excess liquid from the volume defining unit (7).

# EXAMPLE 1. BEADS AS SURFACE ENLARGEMENTS AND CARRIER FOR SINGLE STRANDED DNA HYBRIDISED TO A PRIMER

Synthesis of coating agent (PEG-PEI adduct): 0.43 g of polyethylenimine (Polymin SN from BASF) was dissolved in 45 ml of 50 mM sodium borate buffer (pH 9.5) at 45°C. 5 g of glycidyl ether of monomethoxypolyethylene glycol (Mw 5 000) was added during stirring and the mixture was continuously stirred for 3 h at 45°C.

Surface treatment: A polycarbonate (polycarbonate of bisphenol A. Macrolon DP-1265, Bayer AG, Germany) disc as described above was placed in a plasma reactor (Plasma Science PS0500, BOC Coating Technology, USA) and treated with an oxygen plasma at 5 sccm gas flow and 500 W RF power for 10 min. After venting the reactor, the disc was immersed in a 0.1% solution of the PEG-PEI adduct in borate buffer pH 9.5 for 1 h. The disc was then rinsed with distilled water, blown dry with nitrogen and the water contact angle (sessile drop) was measured on a Ramé-Hart manual goniometer bench. The average of six equilibrium measurements (three droplets) was 24 degrees. An XPS spectrum of the treated surface gave the following molar elemental composition: 73.2%C, 3.7%N, 23.1%O, showing that the surface was essentially covered by the adsorbed

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PEG-PEI adduct.

The microchannel structure was covered with a silicone rubber lid.

Streptavidin-Source 15µm particles: Source 15µm particles were oxidised with periodate, coupled with 6-aminohexanoic acid and further reacted with N-hydroxy-succinimid. Streptavidin (8mg/ml particles) was coupled to the NHS-activated particles at pH 8. Biotin capacity: 0.4 mmol/ml.

### Placing SA-beads in a microchannel structure:

20  $\mu$ l of a 10% Source –SA slurry were added to 0.5  $\mu$ l tube and the beads washed with 1xBW. 20  $\mu$ l BW buffer; 2,5 $\mu$ l double stranded DNA (template hybridised to the primer, (5pmol/ $\mu$ l) and 7.5 $\mu$ l TE and were added, mixed with the beads and incubated at 65°C for 10 min. The beads were then washed in TE once and TE added to a final volume of 20  $\mu$ l. After each step the tube was centrifuged (30 sec.; 10.000 rpm) and the supernatant discarded.

The particles with immobilised DNA were applied as a 2% slurry to a column just before section I (about 8 nl) of the reaction chamber (10) of the microchannel structure described in figure 4.

### Pyrosequencing reaction on the CD device

To minimise the risk for PPi contamination in the pyrosequencing mix, the test tubes used to prepare the mix were washed with 99% EtOH followed by milliQ, and dried upside down overnight.

The Pyrosequencing mix (50 µl) was prepared from the following:

33.5 µl 1xTAE

5 μl Stock A

1 μl 1xTE

4 μl Luciferase (150 ng/μl)

 $2.5 \mu l$  Sulphurylase ( $20 \text{ mU/}\mu l$ )

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#### Stepwise primer extension and detection of nucleotide insertion:

Pyrosequencing mixes with nucleotides ordered in accordance with the template sequence were distributed with intermediate TAE washing to the applicator area (3). Replacement of reagents was accomplished by spinning the disc. The pyrosequencing reaction in the CD device was measured in the detector Ppy1: 1. The reaction chamber functioned as the detection chamber.

Signals were obtained which corresponded to each addition of nucleotide which could be distinguished from the background noise.

# EXAMPLE 2. THE SURFACE OF THE REACTION CHAMBER AS CARRIER OF SINGLE STRANDED DNA HYBRISED TO A PRIMER

Surface treatment and immobilisation of DNA: The surface of each reaction chamber (10) was masked with Owoco Rosa (Owoco AB, Stockholm – Trangsund, Sweden). The structures were then plasma treated as described in example 1 meaning that the unmasked areas were hydrophilized. After removal of Owoco Rose, hydrophobic breaks as indicated above were made by an over-head pen (permanent ink) (Snowman, Japan). The microchannel structures were then covered with a silicone rubber lid and the channels flushed with the PEI-PEG adduct described in example 1, which adhered to the plasma treated surfaces. Thereafter strepavidin was adsorbed (3x) to the surfaces of the reaction chambers followed by a wash with TE. The reaction chamber was then filled with a solution of double stranded DNA (primer DNA hybridised to template DNA, 5pmol/µl) and incubated for 20-30 minutes to immobilise the double-stranded DNA. The channels were then washed twice with TAE.

Stepwise primer extension and detection of nucleotide insertion: See example 1.

3.

## **CLAIMS**

	1.	A method of determining a nucleotide base in a nucleic acid sample comprising the steps of:				
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		(i)	incubating the nucleic acid sample with a primer, DNA polymerase deoxynucleotide triphosphate, or the corresponding			
			deoxynucleotide triphosphate analogue or dideoxynucleotide			
10		<i>(</i> **)	triphosphate (representing a single base?)			
10		(ii)	measuring the pyrophosphate released in step (i)			
	•	(iii)	identifying the nature of the base added by measuring which			
			nucleotide caused the release of PPi in step (ii)			
15		chara	cterised in that steps (i) to (iii) are performed in a microfluidic device.			
13	2.	A method for identifying the sequence of a portion of sample DNA, which method comprises:				
		(i)	forming immobilised double stranded DNA on one or more			
20		(-)	reaction areas in a microchannel structure of a microfluidic device:			
		(ii)	adding a known deoxynucleotide (or the corresponding			
25	,	(11)	deoxynucleotide analogue or dideoxynucleotide) and a DNA			
			polymerase to each of said one or more reaction areas so that			
			•			
			extension of primer only occurs if there is a complementarity of the			
			added deoxynucleotide or dideoxynucleotide with the strand of			
	•		sample DNA that is part of the immobilised double stranded DNA;			
		(iii)	detecting whether or not the deoxynucleotide or dideoxynucleotide			
			added in step (ii) has been added to the primer DNA in said one or			
			more reaction areas,			
30		(iv)	repeating steps (ii) and (iii) as required with a different			
			deoxynucleotide (or the corresponding deoxynucleotide analogue			
			or dideoxynucleotide).			

A method of determining a nucleotide base in a nucleic acid sample

PCT/EP99/10347

according to claim 1 or 2 comprising the steps of:

(i) attaching 0.1 - 200 pmol of a primer or single stranded DNA sample to each of between one and 100,000 pre-determined areas 5 on the surface of a microfluidic device; (ii) hybridising small amounts, e.g. 0.1 - 200 pmol, of single stranded sample DNA or primer respectively to each of the predetermined areas: (iii) adding a known deoxynucleotide, deoxynucleotide analogue or 10 dideoxynucleotide and a DNA polymerase so that extension of the primer only occurs, with consequent release of pyrophosphate (PPi), if there is a complementarity with the sample DNA; (iv) measuring the release of PPi and from which predetermined area on the device it is released; 15 repeating steps (iii) and (iv) as required to construct a DNA (v) sequence for the elongated primers, and hence for portions of the sample DNA. 4. A method for identifying the sequence of a portion of sample DNA, which 20 method comprises: (i) adding sample DNA to a predetermined area on a microfluidic device (ii) moving the sample to a reaction chamber on the microfluidic 25 device (iii) attaching the sample DNA to a surface of the reaction chamber, alternatively hybridising the sample DNA in a single stranded form to a primer attached to the reaction chamber (then to (v)) if the sample DNA has not been attached to a primer attached to the (iv) 30 reaction chamber, hybridising a primer to the DNA in a single stranded form

extending the primer in the presence of a DNA polymerase with a

known deoxynucleotide (dNTP), deoxynucleotide analogue or dideoxynucleotide (ddNTP) such extension being indicated by

(v)

#### WO 00/40750

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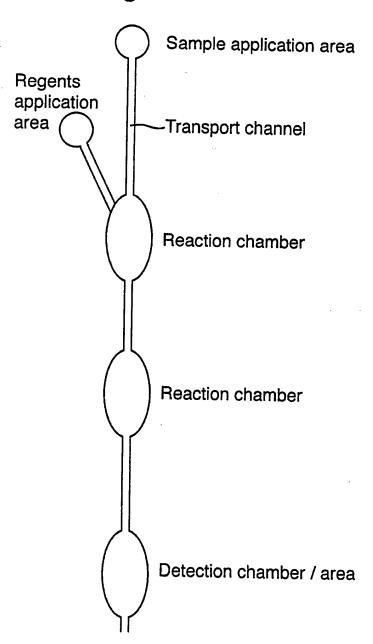


#### PCT/EP99/10347

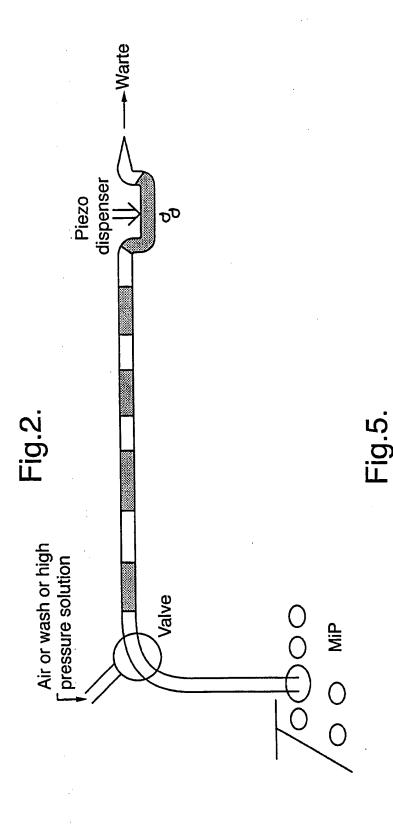
- detection of pyrophosphate (PPi) released from the extension reaction
- (vi) repeating step (v) as required to establish the sequence of the extended primer.
- 5. A method according to any one of claims 1, 3 or 4 wherein the pyrophosphate release is detected by light emitted from a luciferin luciferase reaction.
- 10 6. A method according to claim 2 wherein the detection step involves labelled terminator
  - 7. A method Claim 1-6 wherein the detection of the deoxynucleotide/dideoxynucleotide incorporation is performed in real time.
  - 8. A method according to any one of claims 1-7 wherein microfluidic devices is a disc wherein the fluids maybe moved by centripetal force.



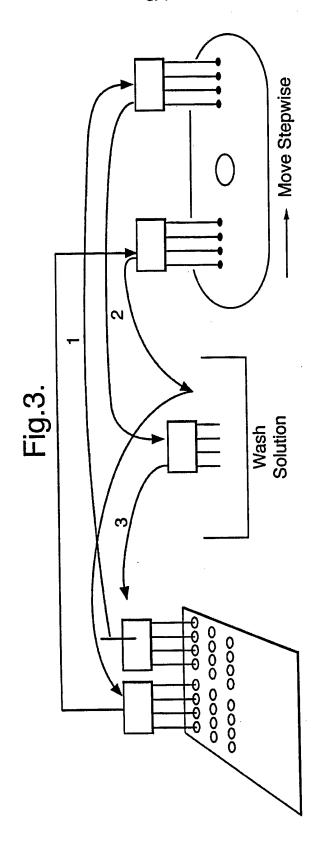
Fig.1.



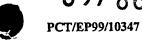
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	=	1	×	×
	=	×	ı	×
	_	ı	ı	ı
	E( mm)	1800	1800	1800
	D( μm)	300	150	300
	C( µm)	300	300	300
	B( mm)	300	300	300
	A( μm)	400	400	400
	L4(µm)	3500	3500	3500
	L3(µm)	4500	4500	4500
	L2(µm)	3500	3000	3000
	L1(µm)	5500	5500	5500
	Channel	K7.K10	K8.K11	K9,K12



SUBSTITUTE SHEET (RULE 26)



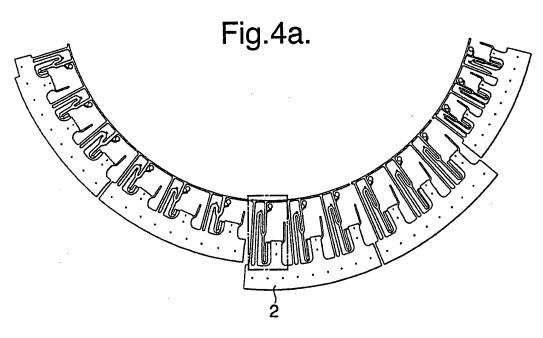


Fig.4b.

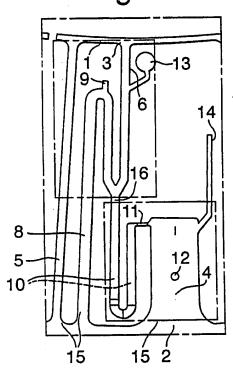


Fig.4c.

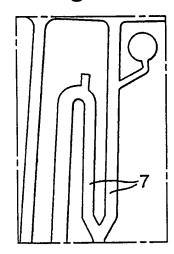


Fig.4d.

